

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/574,625
Source: IFWP
Date Processed by STIC: 4/13/06

ENTERED

CRF Errors Edited by the STIC Systems Branch

Serial Number: 10/574,625

CRF Edit Date: 4/13/06
Edited by: h

— Realigned nucleic acid/amino acid numbers/text in cases where the sequence text "wrapped" to the next line

— Corrected the SEQ ID NO. Sequence numbers edited were:

— Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

  Deleted: invalid beginning/end-of-file text ; page numbers

— Inserted mandatory headings/numeric identifiers, specifically:

— Moved responses to same line as heading/numeric identifier, specifically:

— Other:



IFWP

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/574,625

DATE: 04/13/2006

TIME: 20:17:08

Input Set : A:\PTO.txt
 Output Set: N:\CRF4\04132006\J574625.raw

3 <110> APPLICANT: Hoff, Tine
 5 <120> TITLE OF INVENTION: A Method of Screening For Protein Secreting Recombinant Host
 6 Cells
 8 <130> FILE REFERENCE: 10355.204-US
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/574,625
 C--> 10 <141> CURRENT FILING DATE: 2006-04-04
 10 <160> NUMBER OF SEQ ID NOS: 11
 12 <170> SOFTWARE: PatentIn version 3.3
 14 <210> SEQ ID NO: 1
 15 <211> LENGTH: 2349
 16 <212> TYPE: DNA
 17 <213> ORGANISM: Bacillus subtilis
 20 <220> FEATURE:
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 22 <222> LOCATION: (1000)..(2349)
 23 <223> OTHER INFORMATION: 1-999 is promoter
 25 <220> FEATURE:
 26 <221> NAME/KEY: CDS
 27 <222> LOCATION: (1000)..(2349)
 29 <400> SEQUENCE: 1
 30 tcttcaagg attcatgttt gtttaccaac ctgttctgta agcagttcat attttctcag 60
 32 gtttcttca aatacctcat caaaaacgtc cggcacagag gctgtatca cctcagctcc 120
 34 ctctcccggtt attccgcctt tggttgcac acgttccagc gtctcctcga aagacatatt 180
 36 tttttcgatc agcatcttgc cagttccgtt tagcgaatga atcagaaaat caaaggcttc 240
 38 ttctttggac aggctgtgt ttctgacggc agacagtgc agttcttcaa agattgcagc 300
 40 tatgaatccc ggtccgagc tcgttaaatt gctggccaca tctaaattcg attctttgat 360
 42 ttcccgtaca cggctgaaaa ccgataacaa ttcattcaga cgttctttt tctctgcagc 420
 44 cagtgcttca ctgtgaacga caagttagat gcccgttct gcttcggacg taatggcagg 480
 46 aataaacacgt gagataccgg ctctgtttc tgcctccaaa agacgcagcg gcacacccggc 540
 48 agctatggat acgatgtgag tatttctgtt cacatacggg tacagacggc gcattgtttc 600
 50 gatgacatgg agtggccggaa cgcataatcaa aatcaattgg cacgtatttg cccaaattctc 660
 52 caacggatca gccgataacgt ttggataatc tgacatgagt gcccgcagcc gctccccctt 720
 54 cgttctcgtc tcaataaaata gctcattctc ttttatttgc tcatgtttca acagctgtct 780
 56 agcgatcata tccgccatgc tgccatatcc aatcaatcca atctgttcca tgcactcagt 840
 58 ccttcatata acaatatgaa gtgtaccgtt ttccgcactt tttcacaatt tccccataatc 900
 60 ttttcatttt tatcccacag tttttgttta tgataaaactc aagtcataaa cctatcaata 960
 62 taaatagaca tgtgaaaata gagaaacgga gtgaacatg atg gat aac tat cgt 1014
 63 Met Asp Asn Tyr Arg
 64 1 5
 66 gat gaa aac aga acg aaa ggt aat gag aat gag gtc ttt tta acg aaa 1062
 67 Asp Glu Asn Arg Thr Lys Gly Asn Glu Asn Glu Val Phe Leu Thr Lys
 68 10 15 20
 70 gag aac gat cag agc gcc tcc tac tcg gcc cgc aat gtc att cat gat 1110

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71	Glu	Asn	Asp	Gln	Ser	Ala	Ser	Tyr	Ser	Ala	Arg	Asn	Val	Ile	His	Asp		
72				25					30					35				
74	cag	gag	aag	aaa	aaa	cga	gga	ttc	gga	tgg	ttc	aga	ccg	ttg	ctt	ggc	1158	
75	Gln	Glu	Lys	Lys	Lys	Arg	Gly	Phe	Gly	Trp	Phe	Arg	Pro	Leu	Leu	Gly		
76						40			45			50						
78	gga	gtg	atc	ggc	ggc	agt	ctt	gct	ctt	ggc	att	tac	acg	ttt	aca	ccg	1206	
79	Gly	Val	Ile	Gly	Gly	Ser	Leu	Ala	Leu	Gly	Ile	Tyr	Thr	Phe	Thr	Pro		
80						55			60			65						
82	ctt	ggt	aac	cat	cat	gat	tct	cag	gac	act	gca	aaa	caa	tca	tcc	agc	1254	
83	Leu	Gly	Asn	His	Asp	Ser	Gln	Asp	Thr	Ala	Lys	Gln	Ser	Ser	Ser	Gln		
84	70						75			80			85					
86	cag	caa	acg	caa	tct	gtt	aca	gca	aca	agc	acc	tcc	tct	gaa	tct	aaa	1302	
87	Gln	Gln	Thr	Gln	Ser	Val	Thr	Ala	Thr	Ser	Thr	Ser	Ser	Glu	Ser	Lys		
88						90				95			100					
90	aaa	agc	tca	agc	agc	tca	tct	gca	ttc	aag	agc	gag	gac	tct	tct	aaa	1350	
91	Lys	Ser	Ala	Phe	Lys	Ser	Glu	Asp	Ser	Ser	Lys							
92						105				110			115					
94	atc	tca	gat	atg	gta	gaa	gac	ctt	tca	cca	gcg	att	gtc	ggt	att	aca	1398	
95	Ile	Ser	Asp	Met	Val	Glu	Asp	Leu	Ser	Pro	Ala	Ile	Val	Gly	Ile	Thr		
96						120			125			130						
98	aat	ctt	cag	gca	caa	tca	aac	agc	tct	ttg	ttc	ggc	tct	agt	tct	tct	1446	
99	Asn	Leu	Gln	Ala	Gln	Ser	Asn	Ser	Ser	Leu	Phe	Gly	Ser	Ser	Ser	Ser		
100	135						140				145							
102	gat	tcc	agc	gaa	gat	aca	gaa	agc	ggt	tca	ggg	tca	ggt	gtc	att	ttc	1494	
103	Asp	Ser	Ser	Glu	Asp	Thr	Glu	Ser	Gly	Ser	Gly	Ser	Gly	Val	Ile	Phe		
104	150						155				160			165				
106	aaa	aaa	gag	aat	ggc	aag	gct	tat	atc	att	aca	aat	aac	cac	gtc	gta	1542	
107	Lys	Lys	Glu	Asn	Gly	Lys	Ala	Tyr	Ile	Ile	Thr	Asn	Asn	His	Val	Val		
108						170				175			180					
110	gaa	ggg	gca	tca	tca	ctg	aag	gta	tct	tta	tat	gac	ggc	act	gag	gtt	1590	
111	Glu	Gly	Ala	Ser	Ser	Leu	Lys	Val	Ser	Leu	Tyr	Asp	Gly	Thr	Glu	Val		
112						185			190			195						
114	act	gca	aag	ctg	gta	ggc	agt	gac	tcg	tta	act	gat	tta	gcc	gtc	ctc	1638	
115	Thr	Ala	Lys	Leu	Val	Gly	Ser	Asp	Ser	Leu	Thr	Asp	Leu	Ala	Val	Leu		
116						200			205			210						
118	caa	atc	agt	gat	gac	cac	gtc	aca	aaa	gtg	gca	aac	ttc	ggt	gat	tca	1686	
119	Gln	Ile	Ser	Asp	Asp	His	Val	Thr	Lys	Val	Ala	Asn	Phe	Gly	Asp	Ser		
120	215						220				225							
122	tct	gat	ctt	aga	aca	ggc	gag	acc	gtt	att	gcg	att	ggg	gat	ccg	ctt	1734	
123	Ser	Asp	Leu	Arg	Thr	Gly	Glu	Thr	Val	Ile	Ala	Ile	Gly	Asp	Pro	Leu		
124	230						235				240			245				
126	gga	aaa	gac	ctg	tcc	cgc	aca	gta	aca	caa	gga	att	gta	agc	ggc	gtg	1782	
127	Gly	Lys	Asp	Leu	Ser	Arg	Thr	Val	Thr	Gln	Gly	Ile	Val	Ser	Gly	Val		
128							250			255			260					
130	gac	aga	acg	gtt	tca	atg	tct	aca	tca	gcc	ggc	gaa	acg	agc	att	aac	1830	
131	Asp	Arg	Thr	Val	Ser	Met	Ser	Thr	Ser	Ala	Gly	Glu	Thr	Ser	Ile	Asn		
132						265			270			275						
134	gtc	att	cag	aca	gac	gca	gca	att	aat	cca	ggt	aac	agc	ggc	ggt	cct	1878	
135	Val	Ile	Gln	Thr	Asp	Ala	Ala	Ile	Asn	Pro	Gly	Asn	Ser	Gly	Gly	Pro		

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Input Set : A:\PTO.txt
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136	280	285	290	
138	tgg tta aat aca gac ggc aaa att gtc ggc att aac agt atg aaa atc			1926
139	Leu Leu Asn Thr Asp Gly Lys Ile Val Gly Ile Asn Ser Met Lys Ile			
140	295	300	305	
142	agt gag gat gat gtt gag ggt atc gga ttc gcc att cca agc aat gac			1974
143	Ser Glu Asp Asp Val Glu Gly Ile Gly Phe Ala Ile Pro Ser Asn Asp			
144	310	315	320	325
146	gta aaa ccg att gct gaa gaa ttg ctg tct aaa gga caa att gaa cgt			2022
147	Val Lys Pro Ile Ala Glu Glu Leu Leu Ser Lys Gly Gln Ile Glu Arg			
148	330	335	340	
150	cca tat atc ggt gtc agc atg ctt gat cta gag caa gtg ccg caa aat			2070
151	Pro Tyr Ile Gly Val Ser Met Leu Asp Leu Glu Gln Val Pro Gln Asn			
152	345	350	355	
154	tac caa gaa ggc aca ctc ggc ctg ttc ggc agc cag ctg aat aaa ggc			2118
155	Tyr Gln Glu Gly Thr Leu Gly Leu Phe Gly Ser Gln Leu Asn Lys Gly			
156	360	365	370	
158	gtt tac atc cgt gag gtc gct tca ggc tct cct gct gaa aag gcc gga			2166
159	Val Tyr Ile Arg Glu Val Ala Ser Gly Ser Pro Ala Glu Lys Ala Gly			
160	375	380	385	
162	tta aaa gcg gag gat att atc atc ggc cta aaa ggt aaa gaa att gat			2214
163	Leu Lys Ala Glu Asp Ile Ile Ile Gly Leu Lys Gly Lys Glu Ile Asp			
164	390	395	400	405
166	aca ggc agt gaa ttg cgc aat atc tta tat aaa gac gca aag atc ggt			2262
167	Thr Gly Ser Glu Leu Arg Asn Ile Leu Tyr Lys Asp Ala Lys Ile Gly			
168	410	415	420	
170	gat acc gtt gaa gtg aaa att ctc cga aac ggc aaa gaa atg acg aaa			2310
171	Asp Thr Val Glu Val Lys Ile Leu Arg Asn Gly Lys Glu Met Thr Lys			
172	425	430	435	
174	aaa att aaa ctg gat caa aaa gaa gag aaa act tcg taa			2349
175	Lys Ile Lys Leu Asp Gln Lys Glu Glu Lys Thr Ser			
176	440	445		
179	<210> SEQ ID NO: 2			
180	<211> LENGTH: 449			
181	<212> TYPE: PRT			
182	<213> ORGANISM: Bacillus subtilis			
184	<400> SEQUENCE: 2			
186	Met Asp Asn Tyr Arg Asp Glu Asn Arg Thr Lys Gly Asn Glu Asn Glu			
187	1	5	10	15
190	Val Phe Leu Thr Lys Glu Asn Asp Gln Ser Ala Ser Tyr Ser Ala Arg			
191	20	25	30	
194	Asn Val Ile His Asp Gln Glu Lys Lys Lys Arg Gly Phe Gly Trp Phe			
195	35	40	45	
198	Arg Pro Leu Leu Gly Gly Val Ile Gly Gly Ser Leu Ala Leu Gly Ile			
199	50	55	60	
202	Tyr Thr Phe Thr Pro Leu Gly Asn His Asp Ser Gln Asp Thr Ala Lys			
203	65	70	75	80
206	Gln Ser Ser Ser Gln Gln Thr Gln Ser Val Thr Ala Thr Ser Thr			
207	85	90	95	
210	Ser Ser Glu Ser Lys Ser Ser Ser Ser Ser Ala Phe Lys Ser			

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Input Set : A:\PTO.txt
Output Set: N:\CRF4\04132006\J574625.raw

211 100 105 110
 214 Glu Asp Ser Ser Lys Ile Ser Asp Met Val Glu Asp Leu Ser Pro Ala
 215 115 120 125
 218 Ile Val Gly Ile Thr Asn Leu Gln Ala Gln Ser Asn Ser Ser Leu Phe
 219 130 135 140
 222 Gly Ser Ser Ser Ser Asp Ser Ser Glu Asp Thr Glu Ser Gly Ser Gly
 223 145 150 155 160
 226 Ser Gly Val Ile Phe Lys Lys Glu Asn Gly Lys Ala Tyr Ile Thr
 227 165 170 175
 230 Asn Asn His Val Val Glu Gly Ala Ser Ser Leu Lys Val Ser Leu Tyr
 231 180 185 190
 234 Asp Gly Thr Glu Val Thr Ala Lys Leu Val Gly Ser Asp Ser Leu Thr
 235 195 200 205
 238 Asp Leu Ala Val Leu Gln Ile Ser Asp Asp His Val Thr Lys Val Ala
 239 210 215 220
 242 Asn Phe Gly Asp Ser Ser Asp Leu Arg Thr Gly Glu Thr Val Ile Ala
 243 225 230 235 240
 246 Ile Gly Asp Pro Leu Gly Lys Asp Leu Ser Arg Thr Val Thr Gln Gly
 247 245 250 255
 250 Ile Val Ser Gly Val Asp Arg Thr Val Ser Met Ser Thr Ser Ala Gly
 251 260 265 270
 254 Glu Thr Ser Ile Asn Val Ile Gln Thr Asp Ala Ala Ile Asn Pro Gly
 255 275 280 285
 258 Asn Ser Gly Gly Pro Leu Leu Asn Thr Asp Gly Lys Ile Val Gly Ile
 259 290 295 300
 262 Asn Ser Met Lys Ile Ser Glu Asp Asp Val Glu Gly Ile Gly Phe Ala
 263 305 310 315 320
 266 Ile Pro Ser Asn Asp Val Lys Pro Ile Ala Glu Glu Leu Leu Ser Lys
 267 325 330 335
 270 Gly Gln Ile Glu Arg Pro Tyr Ile Gly Val Ser Met Leu Asp Leu Glu
 271 340 345 350
 274 Gln Val Pro Gln Asn Tyr Gln Glu Gly Thr Leu Gly Leu Phe Gly Ser
 275 355 360 365
 278 Gln Leu Asn Lys Gly Val Tyr Ile Arg Glu Val Ala Ser Gly Ser Pro
 279 370 375 380
 282 Ala Glu Lys Ala Gly Leu Lys Ala Glu Asp Ile Ile Ile Gly Leu Lys
 283 385 390 395 400
 286 Gly Lys Glu Ile Asp Thr Gly Ser Glu Leu Arg Asn Ile Leu Tyr Lys
 287 405 410 415
 290 Asp Ala Lys Ile Gly Asp Thr Val Glu Val Lys Ile Leu Arg Asn Gly
 291 420 425 430
 294 Lys Glu Met Thr Lys Lys Ile Lys Leu Asp Gln Lys Glu Glu Lys Thr
 295 435 440 445
 298 Ser
 302 <210> SEQ ID NO: 3
 303 <211> LENGTH: 8
 304 <212> TYPE: DNA
 305 <213> ORGANISM: Artificial
 307 <220> FEATURE:

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/574,625

DATE: 04/13/2006

TIME: 20:17:08

Input Set : A:\PTO.txt
 Output Set: N:\CRF4\04132006\J574625.raw

308 <223> OTHER INFORMATION: octameric motif
 311 <220> FEATURE:
 W--> 312 <221> NAME/KEY: octameric_motif
 313 <222> LOCATION: (1)..(8)
 315 <400> SEQUENCE: 3
 316 tttcata 8
 319 <210> SEQ ID NO: 4
 320 <211> LENGTH: 44
 321 <212> TYPE: DNA
 322 <213> ORGANISM: Artificial
 324 <220> FEATURE:
 325 <223> OTHER INFORMATION: Primer
 328 <220> FEATURE:
 W--> 329 <221> NAME/KEY: Primer_oth48
 330 <222> LOCATION: (1)..(44)
 332 <400> SEQUENCE: 4 44
 333 gtcatcgat cgcatcggt aatcagacca cttcggtga aggc
 336 <210> SEQ ID NO: 5
 337 <211> LENGTH: 50
 338 <212> TYPE: DNA
 339 <213> ORGANISM: Artificial
 341 <220> FEATURE:
 342 <223> OTHER INFORMATION: Primer
 345 <220> FEATURE:
 W--> 346 <221> NAME/KEY: Primer_oth50
 347 <222> LOCATION: (1)..(50)
 349 <400> SEQUENCE: 5 50
 350 ggagcggatt gaacatgcga ttaaatatcc ttcgagacat tttcgatcgc
 353 <210> SEQ ID NO: 6
 354 <211> LENGTH: 21
 355 <212> TYPE: DNA
 356 <213> ORGANISM: Artificial
 358 <220> FEATURE:
 359 <223> OTHER INFORMATION: Primer
 362 <220> FEATURE:
 W--> 363 <221> NAME/KEY: Primer_260558
 364 <222> LOCATION: (1)..(21)
 366 <400> SEQUENCE: 6
 367 gagtatcgcc agtaaggggc g 21
 370 <210> SEQ ID NO: 7
 371 <211> LENGTH: 44
 372 <212> TYPE: DNA
 373 <213> ORGANISM: Artificial
 375 <220> FEATURE:
 376 <223> OTHER INFORMATION: Primer
 378 <400> SEQUENCE: 7
 379 gccttcaccg gaagtggctt gattagccga tgcgatcgat gaac 44
 382 <210> SEQ ID NO: 8
 383 <211> LENGTH: 23

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/574,625

DATE: 04/13/2006
TIME: 20:17:09

Input Set : A:\PTO.txt
Output Set: N:\CRF4\04132006\J574625.raw

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:3,4,5,6,7,8,9,10,11

VERIFICATION SUMMARY DATE: 04/13/2006
PATENT APPLICATION: US/10/574,625 TIME: 20:17:09

Input Set : A:\PTO.txt
Output Set: N:\CRF4\04132006\J574625.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application No
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:21 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1
L:312 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:3
L:329 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:4
L:346 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:5
L:363 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:6
L:392 M:257 W: Feature value miss-spelled or invalid, <221> Name/Key for SEQ ID#:8
L:409 M:257 W: Feature value miss-spelled or invalid, <221> Name/Key for SEQ ID#:9
L:426 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:10
L:443 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:11

Raw Sequence Listing before editing (for reference only)



IFWP

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/574,625

DATE: 04/13/2006

TIME: 10:17:33

Input Set : A:\01-SQ Listing-04 Apr 2006.txt
 Output Set: N:\CRF4\04132006\J574625.raw

3 <110> APPLICANT: Hoff, Tine
 5 <120> TITLE OF INVENTION: A Method of Screening For Protein Secreting Recombinant Host
 6 Cells
 8 <130> FILE REFERENCE: 10355.204-US
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/574,625
 C--> 10 <141> CURRENT FILING DATE: 2006-04-04
 10 <160> NUMBER OF SEQ ID NOS: 11
 12 <170> SOFTWARE: PatentIn version 3.3

ERRORED SEQUENCES

433 <210> SEQ ID NO: 11
 434 <211> LENGTH: 26
 435 <212> TYPE: DNA
 436 <213> ORGANISM: Artificial
 438 <220> FEATURE:
 439 <223> OTHER INFORMATION: Primer
 442 <220> FEATURE:
 W--> 443 <221> NAME/KEY: Primer_YKDAP1
 444 <222> LOCATION: (1)..(26)
 446 <400> SEQUENCE: 11
 447 gcggatccga tgatgaatga cattgc
 E--> 453 8

Does Not Comply
 Corrected Diskette Needed

26

VERIFICATION SUMMARY

DATE: 04/13/2006

PATENT APPLICATION: US/10/574,625

TIME: 10:17:34

Input Set : A:\01-SQ Listing-04 Apr 2006.txt
Output Set: N:\CRF4\04132006\J574625.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application No
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:21 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1
L:312 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:3
L:329 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:4
L:346 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:5
L:363 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:6
L:392 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:8
L:409 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:9
L:426 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:10
L:443 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:11
L:453 M:254 E: No. of Bases conflict, this line has no nucleotides.